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HH Do, S Melnik, E Rahm - Web, Web-Services, and Database Systems: Node 2002 Web-And 2003 - books.google.com ... siveness thus preventing a qualitative comparison between systems. ... effort and the hardness of match problems. ... very helpful to better compare the effectiveness ...

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O Bachar, D Fischer, R Nussinov, H Wolfson - Protein Engineering Design and Selection, 2003 - Oxford Univ Press ... two typical structure comparison problems and compare them with ... provides an option to allow self comparison, which produces the best match besides die ...

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A Fedorov, AF Merican, W Gilbert - Proceedings of the National Academy of Sciences of the 2002 - National Acad Sciences ... alignment to compare these 30 genes to fungal-intron ... introns from the best fungal-animal match onto the ... such an animal-plant-fungal comparison, and Table ...

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A Krogh, I Brown - J. Mol. Biol., 1994 - cab.zju.edu.cn ... of an HMM are estimated from a training set of unaligned sequences. ... To demonstrate the method, we examine three protein families. Each family consists of a set ...

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MJ Wise - Proceedings of the twenty-seventh SIGCSE technical symposium 1996 - portal.acm.org ... the problem exists of finding the initial set of unique ... to reduce the otherwise $O(n^2$, cost of this comparison. ... of maximal-matches length L do remove match(p, t ...

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WW Cohen, J Richman - Proceedings of the eighth ACM SIGKDD international 2002 - portal.acm.org ... and • in Step 3d, let $Tiht = 0$ (i.e., only remove a from the set of PossibleCenters ... Two of the datasets require clustering, and two require match- ing ...

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F Giunchiglia, P Shvaiko, M Yatskevich - The Semantic Web: Research and Applications: First European 2004 - books.google.com ... step 3) and, finally (step 4), we compare C 2 in ... Evaluation We have done some preliminary comparison between S ... is an estimate of the post match efforts needed ...

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LP Cox, CD Murray, BD Noble - ACM SIGOPS Operating Systems Review, 2002 - portal.acm.org ... meta-data is not chunked and is mu-table. ... with commonly-appearing chunkids, hoping for a false match. ... We call these the add set, delete set, and meta-data ...

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AL Delcher, A Phillips, J Carlton, SL Salzberg - Nucleic Acids Research, 2002 - Oxford Univ Press ... 4 k + 1 bytes for a minimum match size of ... algorithmic extensions that permit alignment and comparison of protein ... It has been used to compare sequences as long ...

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E Brill - Proceedings of the 31st Annual Meeting of the Association ..., 1993 - aci.ldc.upenn.edu
... presented above allow us to **compare** the transformation ... carried out are integer addition
and integer **comparison**. ... hope that this grammar will **match** the grammar ...
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